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301	CGCACGGGGAGCATCTCCTACATCAACATCATGATGCCTTCGGTGTTCGGCACCATCTGC	360
361	CTCCTGGGCATCATCGGGAACTCCACGGTCATCTTCGCGGTCGTGAAGAAGTCCAAGCTG	420
421	CACTGGTGCAACAACGTCCCCGACATCTTCATCATCAACCTCTCGGTAGTAGATCTCCTC	480
481	TTTCTCCTGGGCATGCCCTTCATGATCCACCAGCTCATGGGCAATGGGGTGTGGCACTTT	540
541	GGGGAGACCATGTGCACCCTCATCACGGCCATGGATGCCAATAGTCAGTTCACCAGCACC	009
601	TACATCCTGACCGCCATGGCCATTGACCGCTACCTGGCCACTGTCCACCCCATCTCTTCC	099
661	ACGAAGTTCCGGAAGCCCTCTGTGGCCACCCTGGTGATCTGCCTCCTGTGGGCCCTCTCC	720
721	TTCATCAGCATCACCCCTGTGGGCTGTATGCCAGACTCATCCCCTTCCCAGGAGGTGCA	780
781	GTGGGCTGCGGCATACGCCTGCCCAACCCAGACACTGACCTCTACTGGTTCACCCTGTAC	840
841	CAGTTTTTCCTGGCCTTTGCCCTTTTGTGGTCATCACAGCCGCATACGTGAGGATC	006
901	CTGCAGCGCATGACGTCCTCAGTGGCCCCCGCCTCCCAGCGCAGCATCCGGCTGCGGACA	096
961	AAGAGGGTGACCCGCACAGCCATCGCCATCTGTCTGGTCTTCTTTGTGTGCTGGGCACCC	1020
1021	TACTATGTGCTACAGCTGACCCAGTTGTCCATCAGCCGCCCGACCCTCACCTTTGTCTAC	1080
1081	TTATACAATGCGGCCATCAGCTTGGGCTATGCCAACAGCTGCCTCAACCCCTTTGTGTAC	1140
1141	ATCGTGCTCTGTGAGACGTTCCGCAAACGCTTGGTCCTGTCGGTGAAGCCTGCAGCCCAG	1200
1201	GGGCAGCTTCGCGCTGTCAGCAACGCTCAGACGGCTGACGAGGAGGAGGACAGAAAGCAAA	1260
1261	GGCACCIGA	1269

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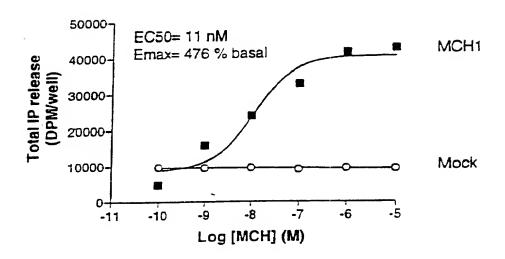
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61	TGCCAGCAACATCTCCGATGGCCAGGATAATCTCACATTGCCGGGGTCACCTCCTCGCAC	120
121	AGGGAGTGTCTCCTACATCAACATCATTATGCCTTCCGTGTTTGGTACCATCTGTCTCT	180
181	GGGCATCGTGGGAAACTCCACGGTCATCTTTGCTGTGGTGAAGAAGTCCAAGCTACACTG	240
241	GTGCAGCAACGTCCCCGACATCTTCATCAACCTCTCTGTGGTGGATCTGCTCTTCCT	300
301	GCTGGGCATGCCTTTCATGATCCACCAGCTCATGGGGAACGGCGTCTGGCACTTTGGGGA	360
361	AACCATGTGCACCCTCATCACAGCCATGGACGCCAACAGTCAGT	420
421	CCTGACTGCCATGACCATTGACCGCTACTTGGCCACCGTCCACCCATCTCCTCCACCAA	480
481	GTTCCGGAAGCCCTCCATGGCCACCCTGGTGATCTGCCTCCTGTGGGCGCTCTCCTTCAT	540
541	CAGTATCACCCCTGTGTGGCTCTACGCCAGGCTCATTCCCTTCCCAGGGGGTGCTGTGGG	009
601	CTGTGGCATCCGCCTGCCAAACCCGGACACTGACCTCTACTGGTTCACTCTGTACCAGTT	099
661	TTTCCTGGCCTTTGCCCTTCCGTTTGTGGTCATTACCGCCGCATACGTGAAAATACTACA	720
721	GCGCATGACGTCTTCGGTGGCCCCAGCCTCCCAACGCAGCATCCGGCTTCGGACAAAGAG	780
781	GGTGACCCGCACGGCCATTGCCATCTGTCTGGTCTTCTTTGTGTGTG	840
941	TGTGCTGCAGCTGACCCAGCTGTCCATCAGCCGCCCGACCCTCACGTTTGTCTACTTGTA	006
901	* CAACGCGGCCATCAGCTTGGGCTATGCTATGCTGCTGCTGAACCCCTTTGTGTATAGT	096
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1081	CIGACAATTCCCCAGTCGCCTCCAAGTCAGGCCACCCCATCAAACCGTGGGGAGAGATAC	1140
1141	TGAGATTAAACCCAAGGCTACCCTGGGAGAATGCAGAGGCTGGAGGCTGGGGGCTTGTAG	1200
1201	CAACCACATTCCAC	1214

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FIGURE 6

IP release in MCH1- and mock-transfected Cos-7 cells



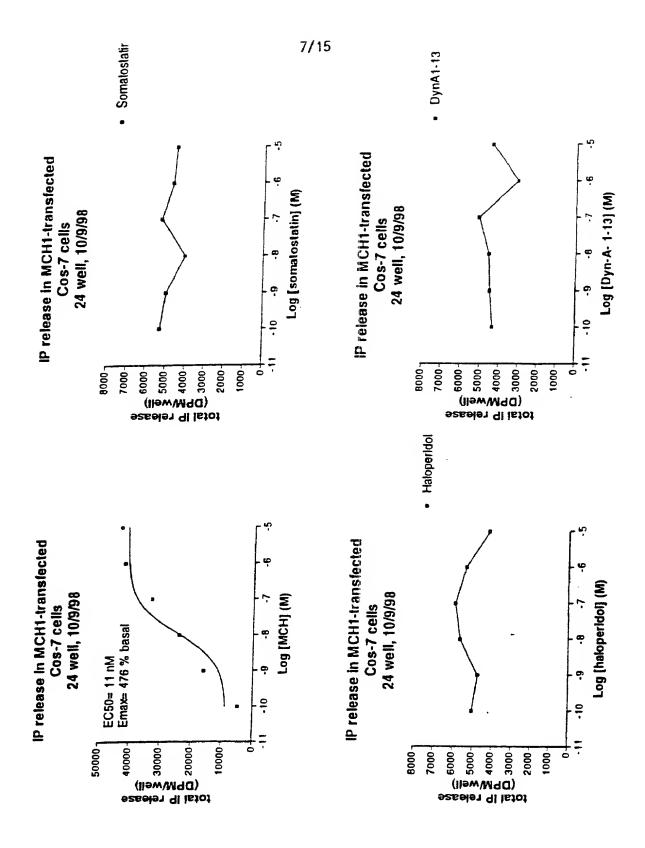
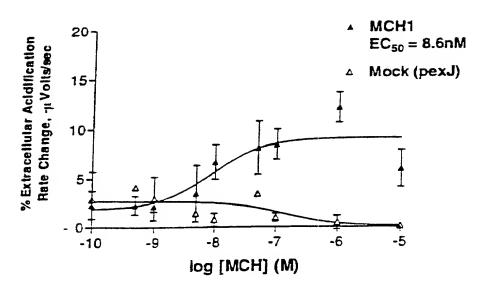


FIGURE 7

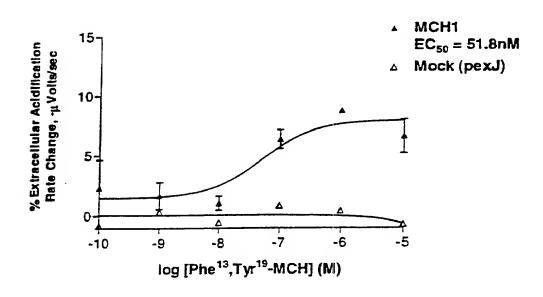
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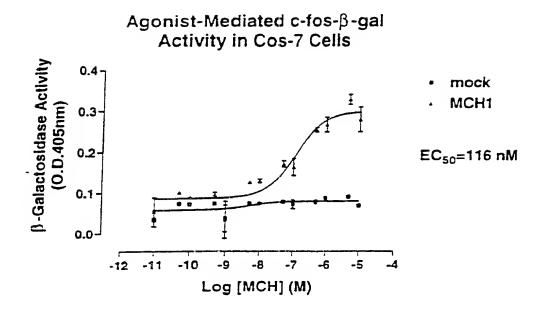
FIGURE

Microphysiometer Response CHO cells



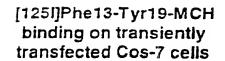
Microphysiometer Response CHO cells

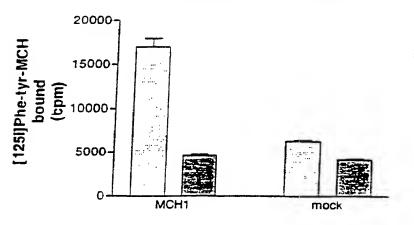




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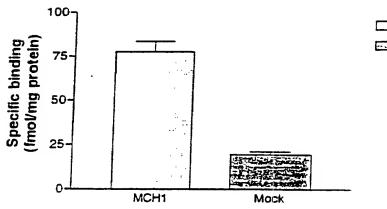
FIGURE 10





☐ Total (cpm)
☐ Non-Specific (cpm)

[125I]Phe13-Tyr19-MCH binding on transiently transfected Cos-7 cells



MCH1
Mock

11/15

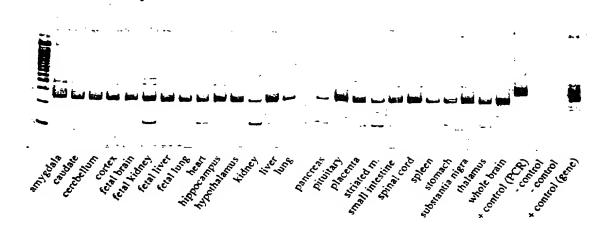


Figure 12

	TL231 R106 R114 B0120	1 MSVGAMKKGV MSVGAMKKGV MSVGAaKKGV	GRAVGLGGGS GRAVGLGGGS GRAVGLGGGS	GCQATEEDPL GCQATEEDPL GCQATEEDPL	PDCGACAPGQ
The state of the s	TL231 R106 R114 B0120	41 GGRRWRLPQP GGRRWRLPQP GGRRWRLPQP	AWVEGSSARL AWVEGSSARL AWVEGSSARL	WEQATGTGWM WEQATGTGWa WEQATGTGWa	80 DLEASLLPTG DLEASLLPTG DLEASLLPTG DLEASLLPTG
At the state of th	TL231 R106 R114 B0120	81 PNASNTSDGP PNASNTSDGP PNASNTSDGP PNASNTSDGP	100 DNLTSAGSPP DNLTSAGSPP DNLTSAGSPP	•	

M S V G A M K K G V G R A V G L G G G S 20 1 D P L P D C G A C A P G Q 40 GCQATEE 21 GGRWRLPQPAWVEGSSARL 60 41 W E Q A T G T G W A D L E A S L L P T G 80 61 PNĀSNTSDGPDNLTSAGSPP 100 81 RTGSISYINIIMPSVFGTIC 120 101 LLGIGNSTVIFAVVKKSKL 140 121 W C N N V P D I F I I N L S V V D L L 160 Н 141 F L L G M P F M I H Q L M G N G V W H F 180 161 GETMCTLITAMDANS QF TST 200 181 VHPISS 220 YILTAMAIDRYLAT 201 TKFRKPSVATLVICLLWALS 240 221 F I S I T P V W L Y A R L I P F P G G A 260 241 DLYWFTLY 280 CGIRLPNPDT V G 261 FLAFALPFVVITAAYVRI 300 QF 281 RMTSSVAPASQRSIRLRT 320 L Q 301 K R V T R T A I A I C L V F 340 VCWAP F 321 Y Y V L Q L T Q L S I S R P 360 T F V Y T L 341 LYNAAISLGYANSCLNPFVY 380 361 IVLCETFRKRLVLSVKPAAQ 400 381 GQLRAVSNAQTADEERTESK 420 401 422 G T 421

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